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Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2008; month=12; day=5; hr=15; min=11; sec=1; ms=756;]

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Reviewer Comments:

<210> 1

<211> 4620

<212> DNA

<213> Homo sapien

Please change the above <213> response to "Homo sapiens". This misspelling also appears in subsequent sequences.

<210> 5

<211> 21

<212> oligonucleotide

<213> artificial sequence

<220>

<223> primers for c-met

<400> 5

agccagtaat gatctcaata g

21

The above <212> response is invalid: use DNA instead. Same error in Sequences 6-8.

Application No: 10599327 Version No: 1.0

Input Set:

Output Set:

Started: 2008-11-17 18:50:41.779
Finished: 2008-11-17 18:50:43.119
Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 340 ms
Total Warnings: 9
Total Errors: 4
No. of SeqIDs Defined: 18
Actual SeqID Count: 18

Error code	Error Description
W 402	Undefined organism found in <213> in SEQ ID (1)
W 402	Undefined organism found in <213> in SEQ ID (2)
E 310	Invalid sequence type in <212> in SEQID: (5)
W 213	Artificial or Unknown found in <213> in SEQ ID (5)
E 310	Invalid sequence type in <212> in SEQID: (6)
W 213	Artificial or Unknown found in <213> in SEQ ID (6)
E 310	Invalid sequence type in <212> in SEQID: (7)
W 213	Artificial or Unknown found in <213> in SEQ ID (7)
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W 402	Undefined organism found in <213> in SEQ ID (15)

SEQUENCE LISTING

<110> Van Andel Research Institute
Shinomiya et al., Nariyoshi

<120> c-met siRNA ADENOVIRUS VECTORS INHIBIT CANCER CELL GROWTH,
INVASION AND TUMORIGENICITY

<130> VAN067 P-328A

<140> 10599327

<141> 2008-11-17

<150> 60/556,473

<151> 2004-03-26

<160> 18

<170> PatentIn version 3.5

<210> 1

<211> 4620

<212> DNA

<213> Homo sapien

<400> 1

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Glu Thr Pro Ile Gln Asn Val Ile Leu His Glu His His Ile Phe Leu
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Pro Cys Gln Asp Cys Ser Ser Lys Ala Asn Leu Ser Gly Gly Val Trp
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Val Ser Ala Leu Gly Ala Lys Val Leu Ser Ser Val Lys Asp Arg Phe		
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His Pro Leu His Ser Ile Ser Val Arg Arg Leu Lys Glu Thr Lys Asp		
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Gly Phe Met Phe Leu Thr Asp Gln Ser Tyr Ile Asp Val Leu Pro Glu		
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245	250	255
Asn Phe Ile Tyr Phe Leu Thr Val Gln Arg Glu Thr Leu Asp Ala Gln		
260	265	270
Thr Phe His Thr Arg Ile Ile Arg Phe Cys Ser Ile Asn Ser Gly Leu		
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His Ser Tyr Met Glu Met Pro Leu Glu Cys Ile Leu Thr Glu Lys Arg		
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Lys Lys Arg Ser Thr Lys Lys Glu Val Phe Asn Ile Leu Gln Ala Ala		
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Tyr Val Ser Lys Pro Gly Ala Gln Leu Ala Arg Gln Ile Gly Ala Ser		
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Leu Asn Asp Asp Ile Leu Phe Gly Val Phe Ala Gln Ser Lys Pro Asp		
340	345	350
Ser Ala Glu Pro Met Asp Arg Ser Ala Met Cys Ala Phe Pro Ile Lys		
355	360	365
Tyr Val Asn Asp Phe Phe Asn Lys Ile Val Asn Lys Asn Asn Val Arg		
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420 425 430

Gln Phe Ser Glu Val Leu Leu Thr Ser Ile Ser Thr Phe Ile Lys Gly
435 440 445

Asp Leu Thr Ile Ala Asn Leu Gly Thr Ser Glu Gly Arg Phe Met Gln
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Leu Asp Ser His Pro Val Ser Pro Glu Val Ile Val Glu His Thr Leu
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Ile Pro Leu Asn Gly Leu Gly Cys Arg His Phe Gln Ser Cys Ser Gln
515 520 525

Cys Leu Ser Ala Pro Pro Phe Val Gln Cys Gly Trp Cys His Asp Lys
530 535 540

Cys Val Arg Ser Glu Glu Cys Leu Ser Gly Thr Trp Thr Gln Gln Ile
545 550 555 560

Cys Leu Pro Ala Ile Tyr Lys Val Phe Pro Asn Ser Ala Pro Leu Glu
565 570 575

Gly Gly Thr Arg Leu Thr Ile Cys Gly Trp Asp Phe Gly Phe Arg Arg
580 585 590

Asn Asn Lys Phe Asp Leu Lys Lys Thr Arg Val Leu Leu Gly Asn Glu
595 600 605

Ser Cys Thr Leu Thr Leu Ser Glu Ser Thr Met Asn Thr Leu Lys Cys

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Pro	Val	Ile	Thr	Ser	Ile	Ser	Pro	Lys	Tyr	Gly	Pro	Met	Ala	Gly	Gly	
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Pro Val Met Ile Ser Met Gly Asn Glu Asn Val Leu Glu Ile Lys Gly
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915 920 925

Trp Lys Gln Ala Ile Ser Ser Thr Val Leu Gly Lys Val Ile Val Gln
930 935 940

Pro Asp Gln Asn Phe Thr Gly Leu Ile Ala Gly Val Val Ser Ile Ser
945 950 955 960

Thr Ala Leu Leu Leu Leu Leu Gly Phe Phe Leu Trp Leu Lys Lys Arg
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Lys Gln Ile Lys Asp Leu Gly Ser Glu Leu Val Arg Tyr Asp Ala Arg
980 985 990

Val His Thr Pro His Leu Asp Arg Leu Val Ser Ala Arg Ser Val Ser
995 1000 1005

Pro Thr Thr Glu Met Val Ser Asn Glu Ser Val Asp Tyr Arg Ala
1010 1015 1020

Thr Phe Pro Glu Asp Gln Phe Pro Asn Ser Ser Gln Asn Gly Ser
1025 1030 1035

Cys Arg Gln Val Gln Tyr Pro Leu Thr Asp Met Ser Pro Ile Leu
1040 1045 1050

Thr Ser Gly Asp Ser Asp Ile Ser Ser Pro Leu Leu Gln Asn Thr
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Asp	Ile	Thr
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